

**Fig. 1**

A

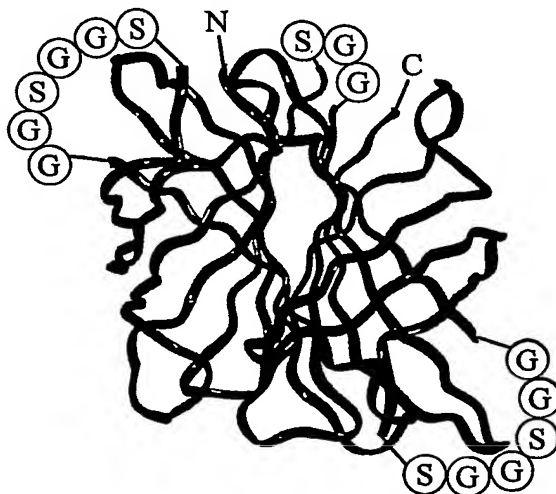
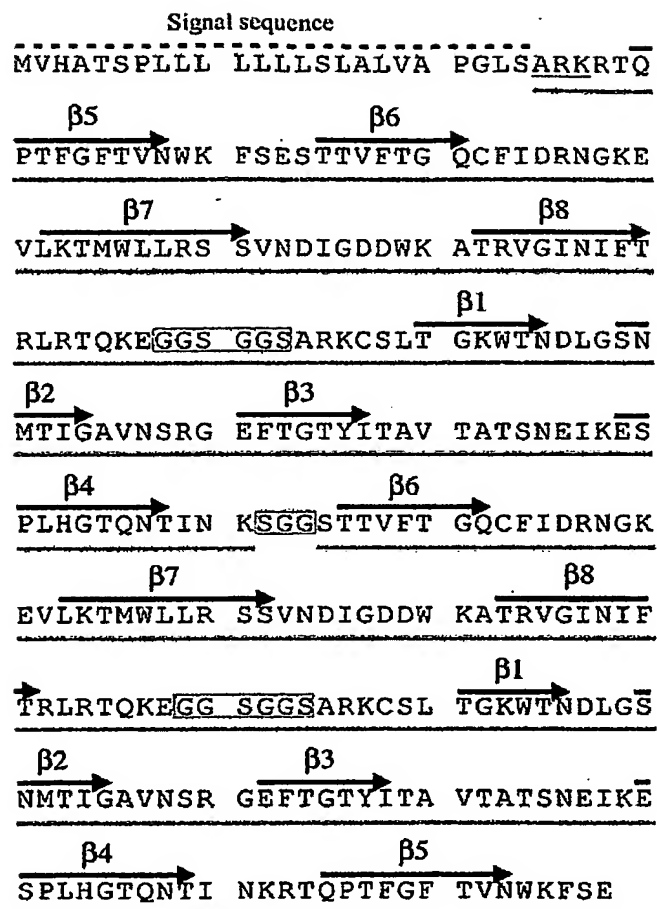
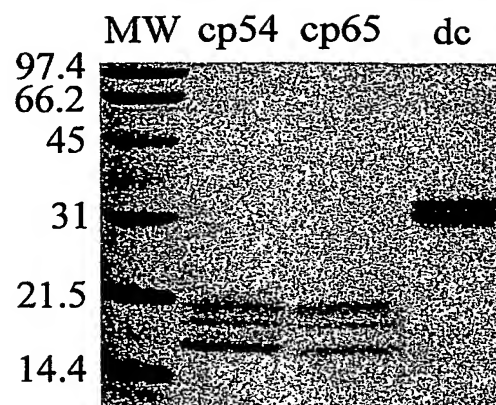
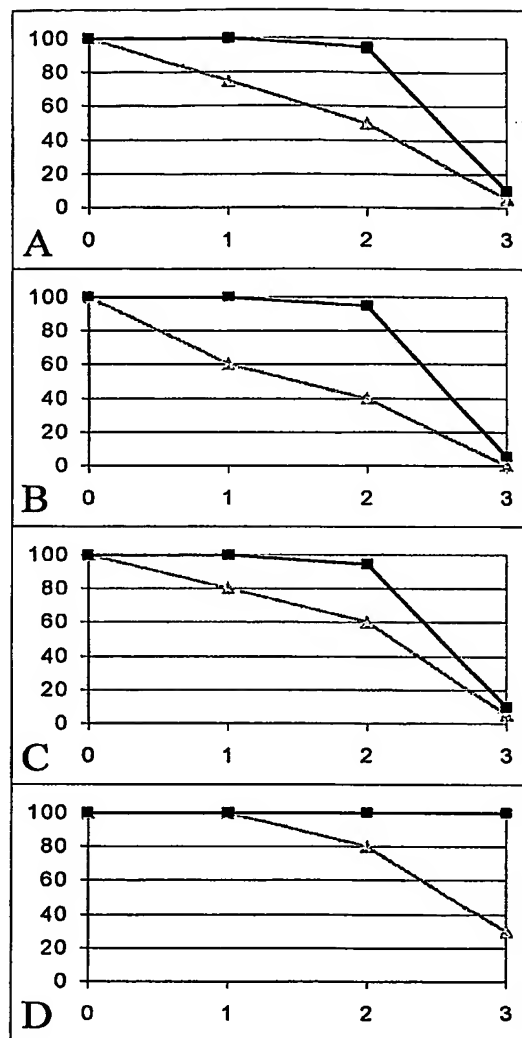


Fig. 2



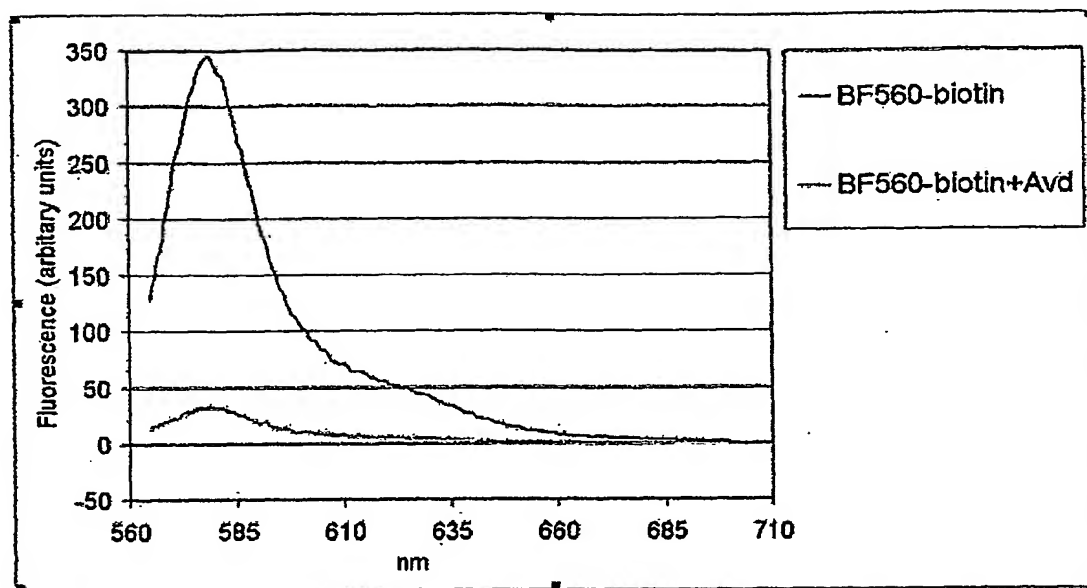
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Fig. 3



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Fig. 4



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Fig. 5

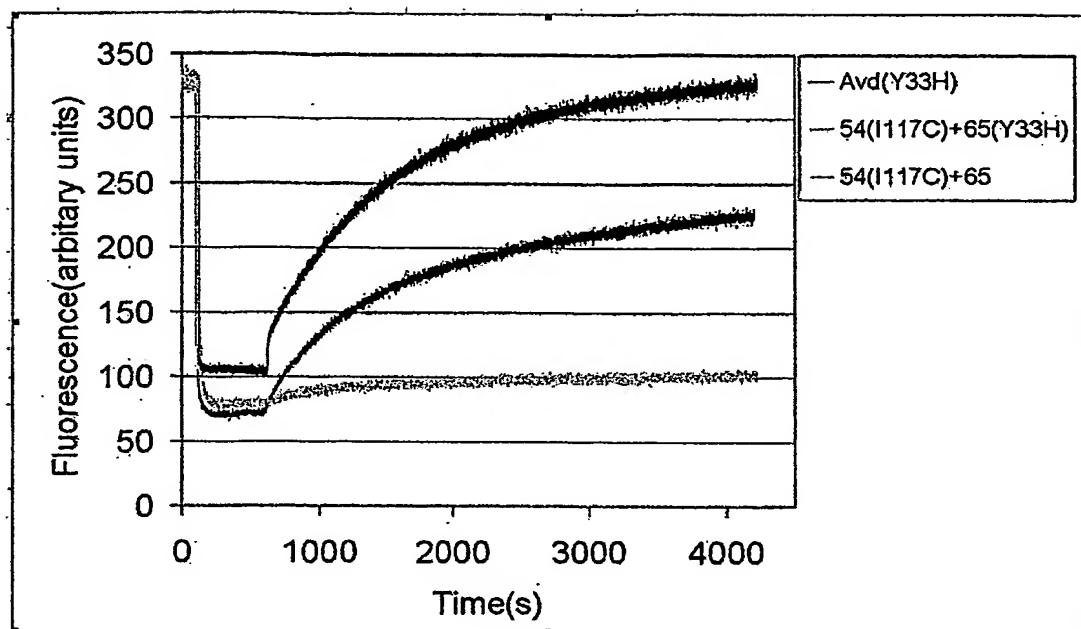


Fig. 6

ATGGTGCACGCAACCTCCCCGCTGCTGCTGCTGCTGCT  
GCTCAGCCTGGCTCTGGTGGCTCCCGGCCTCTCTGCCA  
GGAAGAGGACCCAGCCCACCTTTGGCTTCACCGTCAAT  
TGGAAGTTTTTCAGAGTCCACCACTGTCTTCACGGGCCA  
GTGCTTCATAGACAGGAATGGGAAGGAGGTCCTGAAG  
ACCATGTGGCTGCTGCGGTCAAGTGTTAATGACATTGG  
TGATGACTGGAAAGCTACCAGGGTCGGCATCAACATC  
TTCCTCGCCTGCGCACACAGAAGGAGGGAGGCTCCG  
GAGGCTCCGCCAGAAAGTGCTCGCTGACTGGGAAATG  
GACCAACGATCTGGGCTCCAACATGACCATCGGGGCT  
GTGAACAGCAGAGGTGAATTCACAGGCACCTACATCA  
CAGCCGTAACAGCCACATCAAATGAGATCAAAGAGTC  
ACCACTGCATGGGACACAAAACACCATCAACAAGTCC  
GGCGGATCCACCACTGTCTTCACGGGCCAGTGCTTCAT  
AGACAGGAATGGGAAGGAGGTCCTGAAGACCATGTGG  
CTGCTGCGGTCAAGTGTTAATGACATTGGTGATGACTG  
GAAAGCTACCAGGGTCGGCATCAACATCTTCACTCGCC  
TGCGCACACAGAAGGAGGGAGGCTCCGGAGGCTCCGC  
CAGAAAGTGCTCGCTGACTGGGAAATGGACCAACGAT  
CTGGGCTCCAACATGACCATCGGGGCTGTGAACAGCA  
GAGGTGAATTCACAGGCACCTACATCACAGCCGTAAC  
AGCCACATCAAATGAGATCAAAGAGTCACCACTGCAT  
GGGACACAAAACACCATCAACAAGAGGACCCAGCCCA  
CCTTTGGCTTCACCGTCAATTGGAAGTTTTTCAGAGGGA  
GGTTCCGGATCGGGATCCGGCTCTGGCAGCGGCAGGA  
CCCAGCCCACCTTTGGCTTCACCGTCAATTGGAAGTTT  
TCAGAGTCCACCACTGTCTTCACGGGCCAGTGCTTCAT  
AGACAGGAATGGGAAGGAGGTCCTGAAGACCATGTGG  
CTGCTGCGGTCAAGTGTTAATGACATTGGTGATGACTG  
GAAAGCTACCAGGGTCGGCATCAACATCTTCACTCGCC  
TGCGCACACAGAAGGAGGGAGGCTCCGGAGGCTCCGC  
CAGAAAGTGCTCGCTGACTGGGAAATGGACCAACGAT  
CTGGGCTCCAACATGACCATCGGGGCTGTGAACAGCA  
GAGGTGAATTCACAGGCACCTACATCACAGCCGTAAC  
AGCCACATCAAATGAGATCAAAGAGTCACCACTGCAT  
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AGAAGGAGGGAGGCTCCGGAGGCTCCGCCAGAAAGTG  
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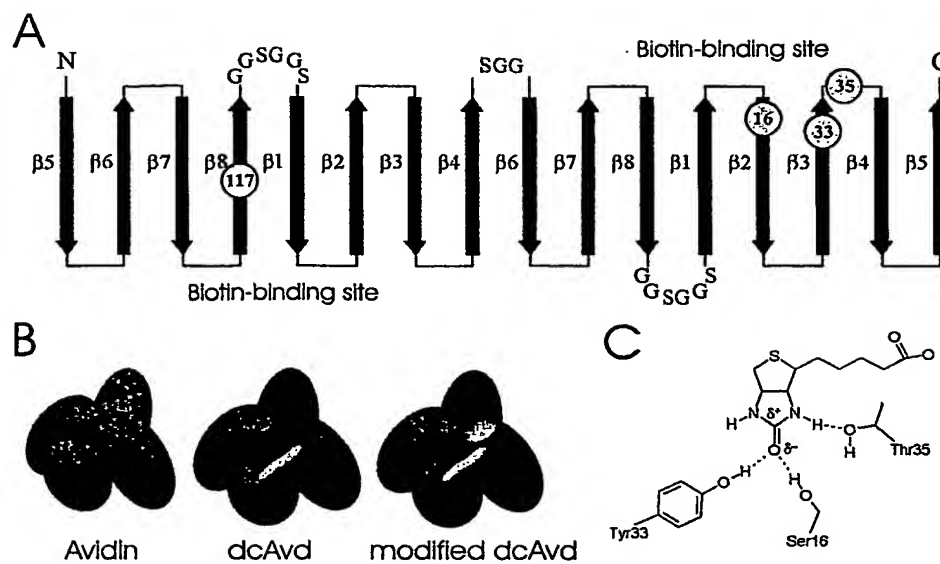
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**Fig. 7**

ATGGTGCACGCAACCTCCCCGCTGCTGCTGCTGCTGCTGCTCA  
GCCTGGCTCTGGTGGCTCCCGGCTCTCTGCCAGGAAGAGGAC  
CCAGCCCACCTTTTGGCTTCACCGTCAATTGGAAGTTTTTCAGAG  
TCCACCCTGTCTTCACGGGCCAGTGCTTCATAGACAGGAATG  
GGAAGGAGGTCCTGAAGACCATGTGGCTGCTGCGGTCAAGTGT  
TAATGACATTGGTGATGACTGGAAAGCTACCAGGGTCGGCATC  
AACATCTTCACTCGCCTGCGCACACAGAAGGAGGGAGGCTCCG  
GAGGCTCCGCCAGAAAGTGCTCGCTGACTGGGAAATGGACCAA  
CGATCTGGGCTCCAACATGACCATCGGGGCTGTGAACAGCAGA  
GGTGAATTCACAGGCACCTACATCACAGCCGTAACAGCCACAT  
CAAATGAGATCAAAGAGTCACCACTGCATGGGACACAAAACAC  
CATCAACAAGTCCGGCGGATCCACCACTGTCTTCACGGGCCAG  
TGCTTCATAGACAGGAATGGGAAGGAGGTCCTGAAGACCATGT  
GGCTGCTGCGGTCAAGTGTTAATGACATTGGTGATGACTGGAA  
AGCTACCAGGGTCGGCATCAACATCTTCACTCGCCTGCGCACA  
CAGAAGGAGGGAGGCTCCGGAGGCTCCGCCAGAAAGTGCTCGC  
TGACTGGGAAATGGACCAACGATCTGGGCTCCAACATGACCAT  
CGGGGCTGTGAACAGCAGAGGTGAATTCACAGGCACCTACATC  
ACAGCCGTAACAGCCACATCAAATGAGATCAAAGAGTCACCAC  
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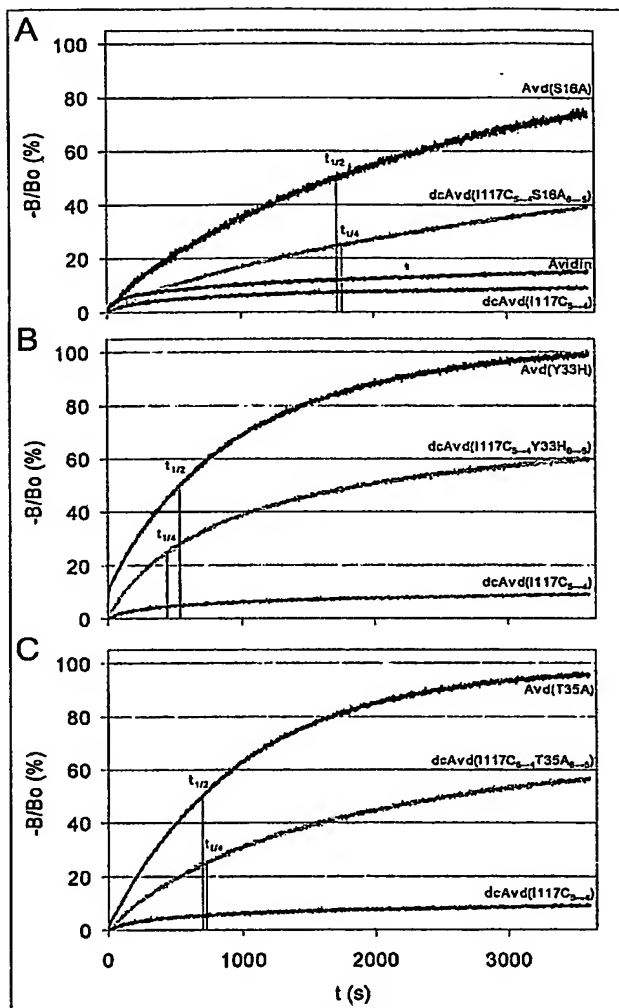
Fig. 8





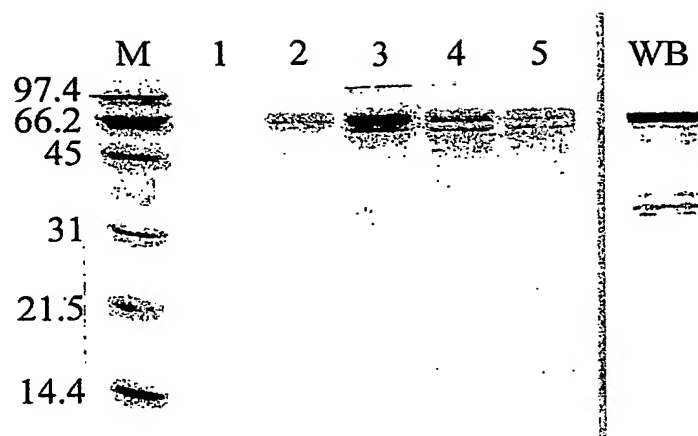
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Fig. 9



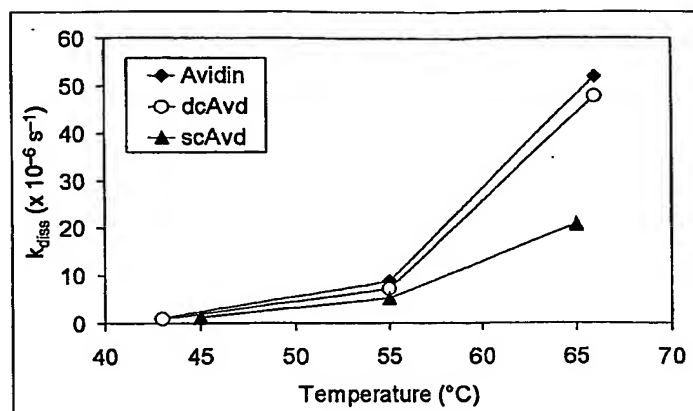
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FIG. 10



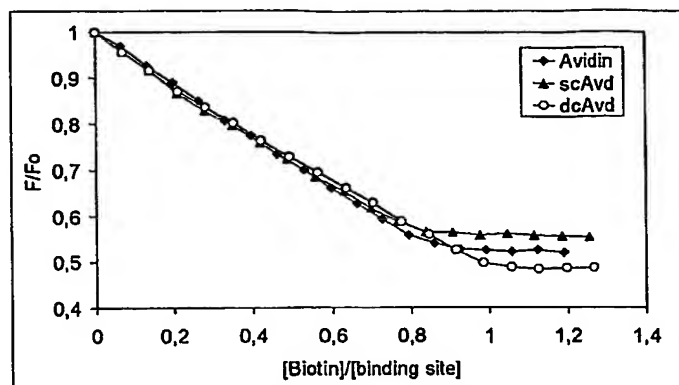
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FIG. 11



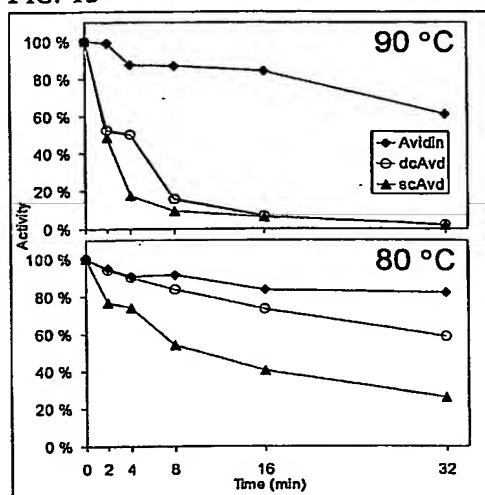
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FIG. 12



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FIG. 13



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Fig. 14

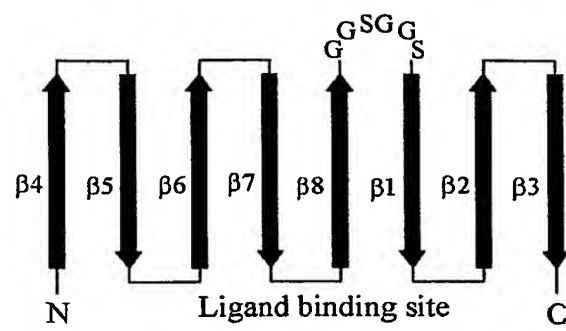
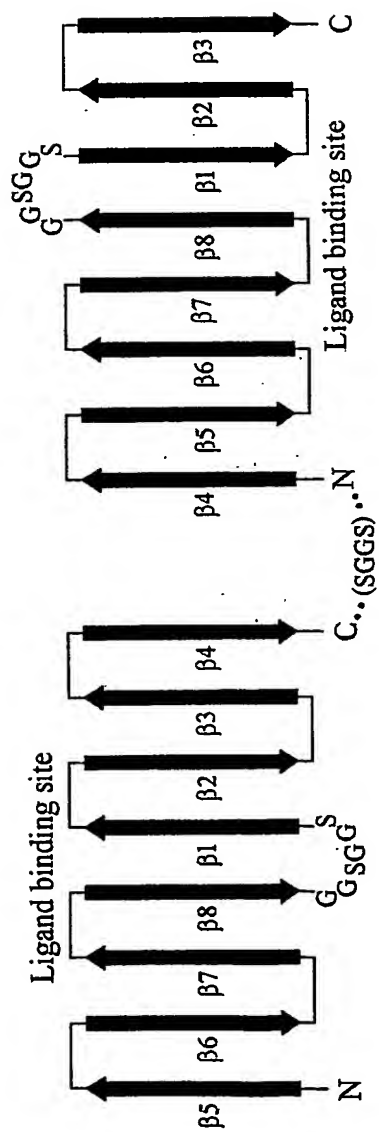
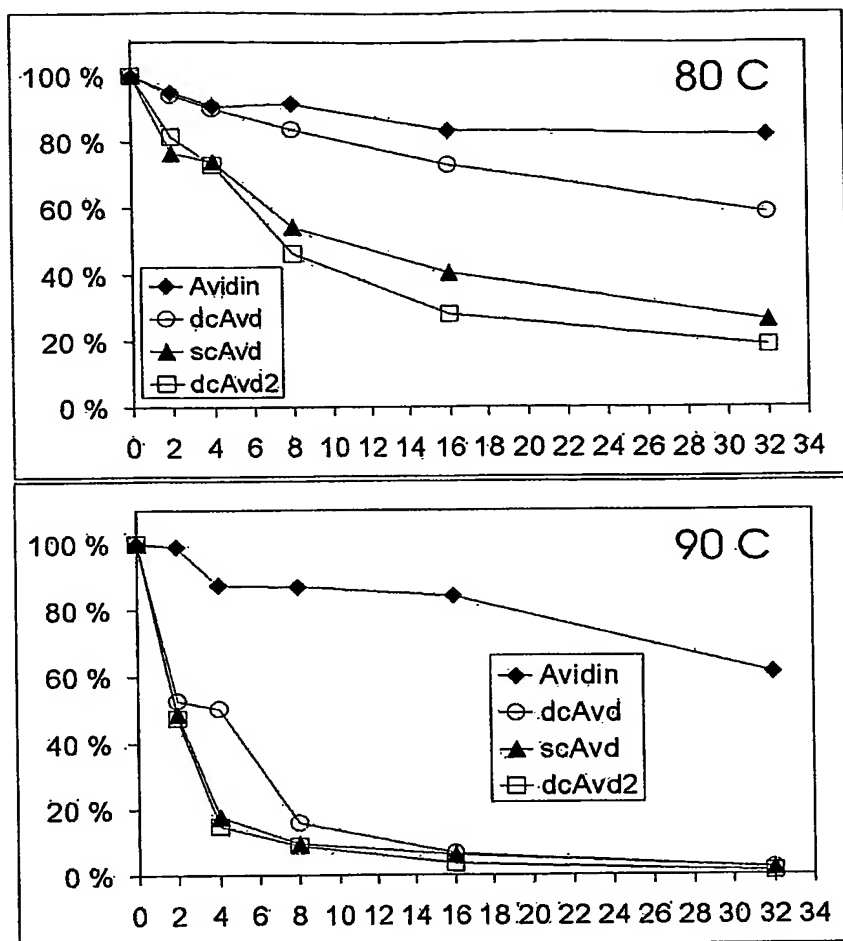


Fig 15.



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Fig. 16





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Fig. 17

1        MNKPSKFALP LAFAAVTASG VASAGTQPTF GFTVNWKFSE  
STTVFTGQCF IDRNGKEVLK  
61        TMWLLRSSVN DIGDDWKATR VGINIFTRLR TQKEGGSGGS  
ARKCSLTGKW TNDLGSNMTI  
121       GAVNSRGEFT GTYITAVTAT SNEIKESPLH GTQNTINKSG  
GSKESPLHGT QNTINKRTQP  
181       TFGFTVNWKF SESTTVFTGQ CFIDRNGKEV LKTMWLLRSS  
VNDIGDDWKA TRVGINIFTR  
241       LRTQKEGGSG GSARKCSLTG KWTNDLGSNM TIGAVNSRGE  
FTGTYITAVT

Fig. 18

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1      ATGAACAAAC CCTCCAAATT CGCTCTGCCG CTTGCCTTCG
CGGCCGTTAC GGCCTCTGGT
61      GTTGCCTCGG CTGGTACCCA GCCCACCTTT GGCTTCACCG
TCAATTGGAA GTTTTCAGAG
121     TCCACCACTG TCTTCACGGG CCAGTGCTTC ATAGACAGGA
ATGGGAAGGA GGTCCCTGAAG
181     ACCATGTGGC TGCTGCGGTC AAGTGTTAAT GACATTGGTG
ATGACTGGAA AGCTACCAGG
241     GTCGGCATCA ACATCTTCAC TCGCCTGCGC ACACAGAAGG
AGGGAGGCTC CGGAGGCTCC
301     GCCAGAAAGT GCTCGCTGAC TGGGAAATGG ACCAACGATC
TGGGCTCCAA CATGACCATC
361     GGGGCTGTGA ACAGCAGAGG TGAATTCACA GGCACCTACA
TCACAGCCGT AACAGCCACA
421     TCAAATGAGA TCAAAGAGTC ACCACTGCAT GGGACACAAA
ACACCATCAA CAAGTCCGGC
481     GGATCCAAAG AGTCACCACT GCATGGGACA CAAAACACCA
TCAACAAGAG GACCCAGCCC
541     ACCTTTGGCT TCACCGTCAA TTGGAAGTTT TCAGAGTCCA
CCACTGTCTT CACGGGCCAG
601     TGCTTCATAG ACAGGAATGG GAAGGAGGTC CTGAAGACCA
TGTGGCTGCT GCGGTCAAGT
661     GTTAATGACA TTGGTGATGA CTGGAAAGCT ACCAGGGTCG
GCATCAACAT CTTCACTCGC
721     CTGCGCACAC AGAAGGAGGG AGGCTCCGGA GGCTCCGCCA
GAAAGTGCTC GCTGACTGGG
781     AAATGGACCA ACGATCTGGG CTCCAACATG ACCATCGGGG
CTGTGAACAG CAGAGGTGAA
841     TTCACAGGCA CCTACATCAC AGCCGTAACA TAA
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